



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/852,238

Source: OIPK

Date Processed by STIC: 08/17/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/852,238

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFT

- 1 ☐ Wrapped Nucleics
Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 ☐ Invalid Line Length
The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 ☐ Misaligned Amino
Numbering
The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 ☐ Non-ASCII
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 ☐ Variable Length
Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 ☐ PatentIn 2.0
"bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s). Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 ☐ Skipped Sequences
(OLD RULES)
Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 ☐ Skipped Sequences
(NEW RULES)
Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 ☐ Use of n's or Xaa's
(NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 ☐ Invalid <213>
Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ☒ Use of <220>
Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 ☐ PatentIn 2.0
"bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 ☐ Misuse of n
n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH - Biotechnology Systems Branch - 08/21/2001

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

BEST AVAILABLE COPY

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/852,238

DATE: 08/17/2001

TIME: 11:40:03

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

3 <110> APPLICANT: Graham P. Allaway et al.
 5 <120> TITLE OF INVENTION: USES OF A CHEMOKINE RECEPTOR FOR INHIBITING HIV-1 INFECTION
 7 <130> FILE REFERENCE: 2048/51320-AB/JPW/SHS
 9 <140> CURRENT APPLICATION NUMBER: 09/852,238
 C--> 10 <141> CURRENT FILING DATE: 2001-05-09
 12 <160> NUMBER OF SEQ ID NOS: 30
 14 <170> SOFTWARE: PatentIn version 3.0
 16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 38
 18 <212> TYPE: DNA
 C--> 19 <213> ORGANISM: artificial
 W--> 21 <220> FEATURE:
 W--> 21 <223> OTHER INFORMATION:
 21 <400> SEQUENCE: 1
 22 caaggctact tccctgattg gcagaactac acaccagg 38
 25 <210> SEQ ID NO: 2
 26 <211> LENGTH: 25
 27 <212> TYPE: DNA
 C--> 28 <213> ORGANISM: artificial
 W--> 30 <220> FEATURE:
 W--> 30 <223> OTHER INFORMATION:
 30 <400> SEQUENCE: 2
 31 agcaagccga gtcctgcgtc gagag 25
 34 <210> SEQ ID NO: 3
 35 <211> LENGTH: 23
 36 <212> TYPE: DNA
 C--> 37 <213> ORGANISM: artificial
 W--> 39 <220> FEATURE:
 W--> 39 <223> OTHER INFORMATION:
 39 <400> SEQUENCE: 3
 40 gggactttcc gctggggact ttc 23
 43 <210> SEQ ID NO: 4
 44 <211> LENGTH: 33
 45 <212> TYPE: DNA
 C--> 46 <213> ORGANISM: artificial
 W--> 48 <220> FEATURE:
 W--> 48 <223> OTHER INFORMATION:
 48 <400> SEQUENCE: 4
 49 cctgttcggg cgccactgct agagattttc cac 33
 52 <210> SEQ ID NO: 5
 53 <211> LENGTH: 31
 54 <212> TYPE: PRT
 55 <213> ORGANISM: human
 57 <400> SEQUENCE: 5
 59 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
 60 1 5 10 15
 62 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg

Does Not Comply
Corrected Diskette Needed

Errored
 When field 213 is artificial or unknown,
 descriptions are required in fields 220
 through 223

BEST AVAILABLE COPY

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/852,238

DATE: 08/17/2001

TIME: 11:40:03

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

```

63          20          25          30
65 <210> SEQ ID NO: 6
66 <211> LENGTH: 15
67 <212> TYPE: PRT
68 <213> ORGANISM: human
70 <400> SEQUENCE: 6
72 His Tyr Ala Ala Ala Gln Trp Asp Phe Gly Asn Thr Met Cys Gln
73 1          5          10          15
75 <210> SEQ ID NO: 7
76 <211> LENGTH: 32
77 <212> TYPE: PRT
78 <213> ORGANISM: human
80 <400> SEQUENCE: 7
82 Arg Ser Gln Lys Glu Gly Leu His Tyr Thr Cys Ser Ser His Phe Pro
83 1          5          10          15
85 Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile Val
86          20          25          30
88 <210> SEQ ID NO: 8
89 <211> LENGTH: 17
90 <212> TYPE: PRT
91 <213> ORGANISM: human
93 <400> SEQUENCE: 8
95 Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser Asn Arg Leu Asp
96 1          5          10          15
98 Gln
101 <210> SEQ ID NO: 9
102 <211> LENGTH: 36
103 <212> TYPE: DNA
C--> 104 <213> ORGANISM: artificial
W--> 106 <220> FEATURE:
W--> 106 <223> OTHER INFORMATION:
106 <400> SEQUENCE: 9
107 aagcttggag aaccagcggt taccatggag gggatc 36
110 <210> SEQ ID NO: 10
111 <211> LENGTH: 30
112 <212> TYPE: DNA
C--> 113 <213> ORGANISM: artificial
W--> 115 <220> FEATURE:
W--> 115 <223> OTHER INFORMATION:
115 <400> SEQUENCE: 10
116 gtctgagtct gagtcaagct tggagaacca 30
119 <210> SEQ ID NO: 11
120 <211> LENGTH: 41
121 <212> TYPE: DNA
C--> 122 <213> ORGANISM: artificial
W--> 124 <220> FEATURE:
W--> 124 <223> OTHER INFORMATION:
124 <400> SEQUENCE: 11
125 ctcgagcatc tgtgttagct ggagtgaaaa cttgaagact c 41

```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/852,238

DATE: 08/17/2001

TIME: 11:40:03

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

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128 <210> SEQ ID NO: 12
129 <211> LENGTH: 30
130 <212> TYPE: DNA
C--> 131 <213> ORGANISM: artificial
W--> 133 <220> FEATURE:
W--> 133 <223> OTHER INFORMATION:
133 <400> SEQUENCE: 12
134 gtctgagtct gagtcctcga gcatctgtgt      30
137 <210> SEQ ID NO: 13
138 <211> LENGTH: 32
139 <212> TYPE: DNA
C--> 140 <213> ORGANISM: artificial
W--> 142 <220> FEATURE:
W--> 142 <223> OTHER INFORMATION:
142 <400> SEQUENCE: 13
143 aagcttcaga gagaagccgg gatggaaact cc      32
146 <210> SEQ ID NO: 14
147 <211> LENGTH: 30
148 <212> TYPE: DNA
C--> 149 <213> ORGANISM: artificial
W--> 151 <220> FEATURE:
W--> 151 <223> OTHER INFORMATION:
151 <400> SEQUENCE: 14
152 gtctgagtct gagtcaagct tcagagagaa      30
155 <210> SEQ ID NO: 15
156 <211> LENGTH: 32
157 <212> TYPE: DNA
C--> 158 <213> ORGANISM: artificial
W--> 160 <220> FEATURE:
W--> 160 <223> OTHER INFORMATION:
160 <400> SEQUENCE: 15
161 ctcgagctga gtcagaaccc agcagagagt tc      32
164 <210> SEQ ID NO: 16
165 <211> LENGTH: 30
166 <212> TYPE: DNA
C--> 167 <213> ORGANISM: artificial
W--> 169 <220> FEATURE:
W--> 169 <223> OTHER INFORMATION:
169 <400> SEQUENCE: 16
170 gtctgagtct gagtcctcga gctgagtcag      30
173 <210> SEQ ID NO: 17
174 <211> LENGTH: 32
175 <212> TYPE: DNA
C--> 176 <213> ORGANISM: artificial
W--> 178 <220> FEATURE:
W--> 178 <223> OTHER INFORMATION:
178 <400> SEQUENCE: 17
179 aagcttcagt acatccacaa catgctgtcc ac      32
182 <210> SEQ ID NO: 18

```

RAW SEQUENCE LISTING

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TIME: 11:40:03

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

```

183 <211> LENGTH: 30
184 <212> TYPE: DNA
C--> 185 <213> ORGANISM: artificial
W--> 187 <220> FEATURE:
W--> 187 <223> OTHER INFORMATION:
187 <400> SEQUENCE: 18
188 gtctgagtct gagtcaagct tcagtacatc 30
191 <210> SEQ ID NO: 19
192 <211> LENGTH: 31
193 <212> TYPE: DNA
C--> 194 <213> ORGANISM: artificial
W--> 196 <220> FEATURE:
W--> 196 <223> OTHER INFORMATION:
196 <400> SEQUENCE: 19
197 ctcgagcctc gttttataaa ccagccgaga c 31
200 <210> SEQ ID NO: 20
201 <211> LENGTH: 30
202 <212> TYPE: DNA
C--> 203 <213> ORGANISM: artificial
W--> 205 <220> FEATURE:
W--> 205 <223> OTHER INFORMATION:
205 <400> SEQUENCE: 20
206 gtctgagtct gagtccctcga gcctcgtttt 30
209 <210> SEQ ID NO: 21
210 <211> LENGTH: 29
211 <212> TYPE: DNA
C--> 212 <213> ORGANISM: artificial
W--> 214 <220> FEATURE:
W--> 214 <223> OTHER INFORMATION:
214 <400> SEQUENCE: 21
215 aagcttcagg gagaagtgaa atgacaacc 29
218 <210> SEQ ID NO: 22
219 <211> LENGTH: 30
220 <212> TYPE: DNA
C--> 221 <213> ORGANISM: artificial
W--> 223 <220> FEATURE:
W--> 223 <223> OTHER INFORMATION:
223 <400> SEQUENCE: 22
224 gtctgagtct gagtcaagct tcagggagaa 30
227 <210> SEQ ID NO: 23
228 <211> LENGTH: 33
229 <212> TYPE: DNA
C--> 230 <213> ORGANISM: artificial
W--> 232 <220> FEATURE:
W--> 232 <223> OTHER INFORMATION:
232 <400> SEQUENCE: 23
233 ctcgagcaga cctaaaacac aatagagagt tcc 33
236 <210> SEQ ID NO: 24
237 <211> LENGTH: 30

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RAW SEQUENCE LISTING

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DATE: 08/17/2001

TIME: 11:40:03

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

```

238 <212> TYPE: DNA
C--> 239 <213> ORGANISM: artificial
W--> 241 <220> FEATURE:
W--> 241 <223> OTHER INFORMATION:
241 <400> SEQUENCE: 24
242 gtctgagtct gagtccctcga gcagacctaa 30
245 <210> SEQ ID NO: 25
246 <211> LENGTH: 34
247 <212> TYPE: DNA
C--> 248 <213> ORGANISM: artificial
W--> 250 <220> FEATURE:
W--> 250 <223> OTHER INFORMATION:
250 <400> SEQUENCE: 25
251 aagcttctgt agagttaaaa aatgaacccc acgg 34
254 <210> SEQ ID NO: 26
255 <211> LENGTH: 30
256 <212> TYPE: DNA
C--> 257 <213> ORGANISM: artificial
W--> 259 <220> FEATURE:
W--> 259 <223> OTHER INFORMATION:
259 <400> SEQUENCE: 26
260 gtctgagtct gagtcaagct tctgtagagt 30
263 <210> SEQ ID NO: 27
264 <211> LENGTH: 34
265 <212> TYPE: DNA
C--> 266 <213> ORGANISM: artificial
W--> 268 <220> FEATURE:
W--> 268 <223> OTHER INFORMATION:
268 <400> SEQUENCE: 27
269 ctcgagccat ttcatttttc tacaggacag catc 34
272 <210> SEQ ID NO: 28
273 <211> LENGTH: 30
274 <212> TYPE: DNA
C--> 275 <213> ORGANISM: artificial
W--> 277 <220> FEATURE:
W--> 277 <223> OTHER INFORMATION:
277 <400> SEQUENCE: 28
278 gtctgagtct gagtccctcga gccatttcat 30
281 <210> SEQ ID NO: 29
282 <211> LENGTH: 39
283 <212> TYPE: DNA
C--> 284 <213> ORGANISM: artificial
W--> 286 <220> FEATURE:
W--> 286 <223> OTHER INFORMATION:
286 <400> SEQUENCE: 29
287 gtctgagtct gagtcaagct taacaagatg gattatcaa 39
290 <210> SEQ ID NO: 30
291 <211> LENGTH: 39
292 <212> TYPE: DNA

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/852,238

DATE: 08/17/2001

TIME: 11:40:04

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:19 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:21 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:21 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:28 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:30 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:30 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:37 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:39 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:39 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:46 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:48 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:48 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:104 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:106 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:106 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:113 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:115 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:115 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:122 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:124 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:124 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:131 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:133 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:133 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:140 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:142 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:142 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:149 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:151 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:151 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:158 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:160 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:160 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:167 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:169 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:169 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:176 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:178 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:178 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:185 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:187 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:187 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:194 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:196 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:196 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:203 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20
L:205 M:258 W: Mandatory Feature missing, <220> FEATURE:

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/852,238

DATE: 08/17/2001

TIME: 11:40:04

Input Set : A:\51320-AB.txt

Output Set: N:\CRF3\08162001\I852238.raw

L:205 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:212 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
L:214 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:214 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:221 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22
L:223 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:223 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:230 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23
L:232 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:232 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:239 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:241 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:241 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:248 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:250 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:250 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:257 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
L:259 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:259 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:266 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
L:268 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:268 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:275 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28
L:277 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:277 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:284 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:286 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:286 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:293 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30